AMENDMENTS In the Claims

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10.(currently amended) A composition comprising a polymerizing agent polymerase including a molecular tag covalently bonded to a site on the polymerizing agent polymerase and a monomer including nucleotide types for the polymerase, where at least one nucleotide type includes a molecular tag bonded to a part of the nucleotide that is released upon monomer nucleotide incorporation, where at least one of the tags has a fluorescence property that undergoes a change before, during and/or after each of a sequence of monomer nucleotide incorporations due to an interaction between the polymerizing agent polymerase and the monomer nucleotide and where the polymerizing agent polymerase lacks the ability to remove a previously incorporated monomer nucleotide.

11.(canceled)

12.(canceled)

13.(canceled)

14.(canceled)

15.(canceled)

1 16.(currently amended) The composition of claim 10, wherein each of the monomers
2 nucleotide types comprises a deoxynucleotide triphosphate (dNTP) and the monomer nucleotide tag
3 is covalently bonded either directly or through a linker to the pyrophosphate moiety of each its
4 dNTP.

17.(currently amended) The composition of claim 10, wherein the at least one tag comprises
2 a fluorescent tag and the fluorescence property comprises a duration, an intensity and/or frequency
3 of emitted fluorescent light.

18.(currently amended) The composition of claim 1710, wherein the polymerase tag comprises

- a fluorescent tag and wherein the fluorescence property is fluorescence resonance energy transfer

 (FRET), where either the monomer nucleotide tag or the polymerase tag comprises a donor and the

 other tag comprises an acceptor and where FRET occurs when the two tags are in close proximity.
- 1 19.(currently amended) The composition of claim 13108, wherein the polymerase comprises
 2 Taq DNA polymerase I having a tag attached to an amino acid at a specific amino acid position of
 3 the Taq DNA polymerase I, where the amino acid position and the site is or the sites are selected
 4 from the group consisting of 513-518, 643, 647, 649 and 653-661 of SEQ. ID No. 11, where the tag
 5 comprises a fluorescent molecule.
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 - 48.(canceled)
 - 49.(canceled)
 - 1 50.(currently amended)

A composition comprising a polymerizing agent polymerase including

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- a molecular tag covalently bonded to a site on the polymerizing agent polymerase and a 2 deoxynucleotide triphosphate (dNTP) types for the polymerase, where at least one dNTP type 3 including includes a molecular tag covalently bonded directly or through a linker to the 4 pyrophosphate moiety a part of the dNTP the is release upon dNTP incorporation, where at least one 5 of the tags has a fluorescence property that undergoes a change before, during and/or after each of 6 a sequence of monomer dNTP incorporations due to an interaction between the polymerizing agent 7 polymerase and the dNTP.
- The composition of claim 50, wherein the polymerizing agent 51.(currently amended) 1 polymerase is a polymerase or reverse transcriptase. 2
- The composition of claim 5150, wherein the polymerase is selected 52.(currently amended) 1 from the group consisting of Taq DNA polymerase I, T7 DNA polymerase, Sequenase, and the 2 Klenow fragment from E. coli DNA polymerase I. 3
- The composition of claim 51, wherein the reverse transcriptase 53.(previously presented) 1 comprises HIV-1 reverse transcriptase. 2
- The composition of claim 50, wherein at least one of the tags 54.(currently amended) 1 comprises a fluorescent tag and the fluorescence property comprises a duration, an intensity and/or 2 frequency of emitted fluorescent light. 3
- The composition of claim 5450, wherein the polymerase tag comprises 55 (currently amended) 1 a fluorescent tag and wherein the fluorescence property is fluorescence resonance energy transfer 2 (FRET), where either the monomer nucleotide tag or the polymerase tag comprises a donor and the 3 other tag comprises an acceptor and where FRET occurs when the two tags are in close proximity. 4
- The composition of claim 52109, wherein the polymerase comprises 56.(previously presented) 5 Taq DNA polymerase I having a tag attached to an amino acid at a specific amino acid position of 6 the Tay DNA polymerase I, where the amino acid position and the site is or the sites are selected 7 from the group consisting of 513-518, 643, 647, 649 and 653-661 of SEQ. ID No. 11, where the tag 8 comprises a fluorescent molecule. 9

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62.0	(canceled)
63.	canceled)

- 64.(currently amended) A composition comprising a polymerizing agent polymerase including a molecular tag covalently bonded to a site on the polymerizing agent polymerase and a deoxynucleotide triphosphate (dNTP) types for the polymerase, where at least one dNTP type including includes a molecular tag covalently bonded directly or through a linker to the γ phosphate group of the dNTP, where at least one of the tags has a fluorescence property that undergoes a change before, during and/or after each of a sequence of monomer dNTP incorporations due to an interaction between the polymerizing agent polymerase and the dNTP.
- 1 65.(currently amended) The composition of claim 64, wherein the polymerizing agent is a polymerase or comprises a reverse transcriptase.
- 1 66.(currently amended) The composition of claim 6564, wherein the polymerase is selected 2 from the group consisting of Taq DNA polymerase I, T7 DNA polymerase, Sequenase, and the 3 Klenow fragment from E. coli DNA polymerase I.
- 1 67.(previously presented) The composition of claim 65, wherein the reverse transcriptase comprises HIV-1 reverse transcriptase.
- 1 68.(currently amended) The composition of claim 64, wherein at least one of the tags
 2 comprises a fluorescent tag and the fluorescence property comprises a duration, an intensity and/or
 3 frequency of emitted fluorescent light.
- 1 69.(currently amended) The composition of claim 6864, wherein the polymerase tag comprises
 2 a fluorescent tag and wherein the fluorescence property is fluorescence resonance energy transfer
 3 (FRET), where either the monomer nucleotide tag or the polymerase tag comprises a donor and the
 4 other tag comprises an acceptor and where FRET occurs when the two tags are in close proximity.

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_	70.(currently amended)	The composition of claim 66110, wherein the polymerase comprises
5	70.(entrently amended)	and the second position of
6	Taq DNA polymerase I having a tag attached to an amino acid at a specific amino acid position of	
7 .	the Taq DNA polymerase I,	where the amino acid position and the site is or the sites are selected
8	from the group consisting of	513-518, 643, 647, 649 and 653-661 of SEQ. ID No. 11, where the tag
9	comprises a fluorescent mol	ecule.

- A composition comprising a polymerizing agent polymerase including 71.(currently amended) a molecular tag covalently bonded to a site on the polymerizing agent polymerase and a monomer including nucleotide types for the polymerase, where at least one nucleotide type includes a molecular tag covalently bonded directly or through a linker to the terminal phosphate of the monomer nucleotide, where at least one of the tags has a fluorescence property that undergoes a change before, during and/or after each of a sequence of monomer nucleotide incorporations due to an interaction between the polymerizing agent polymerase and the monomer nucleotide.
- The composition of claim 71, wherein the polymerizing agent is a 72.(currently amended) 1 polymerase or comprises a reverse transcriptase. 2
- The composition of claim 7271, wherein the polymerase is selected 73.(currently amended) 1 from the group consisting of Taq DNA polymerase I, T7 DNA polymerase, Sequenase, and the 2 Klenow fragment from E. coli DNA polymerase I. 3
- The composition of claim 72, wherein the reverse transcriptase 74.(previously presented) 1 comprises HIV-1 reverse transcriptase. 2

75.(canceled)

- The composition of claim 71, wherein at least one of the tags 76.(currently amended) 1 comprises a fluorescent tag and the fluorescence property comprises a duration, an intensity and/or 2 frequency of emitted fluorescent light. 3
- The composition of claim 7671, wherein the polymerase tag comprises 77.(currently amended) 1 a fluorescent tag and wherein the fluorescence property is fluorescence resonance energy transfer 2 (FRET), where either the monomer nucleotide tag or the polymerase tag comprises a donor and the 3

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- other tag comprises an acceptor and where FRET occurs when the two tags are in close proximity. 4
- The composition of claim 73111, wherein the polymerase comprises 78.(currently amended) 5
- Taq DNA polymerase I having a tag attached to an amino acid at a specific amino acid position of 6
- the Taq DNA polymerase I, where the amino acid position and the site is or sites are selected from 7
- the group consisting of 513-518, 643, 647, 649 and 653-661 of SEQ. ID No. 11, where the tag 8
- comprises a fluorescent molecule. 9
- A composition comprising a polymerizing agent polymerase including 79.(currently amended) 1
- a molecular tag covalently bonded to a site on the polymerizing agent polymerase lacking 3' to 5' 2
- exonuclease activity and a monomer including nucleotide types for the polymerase, where at least 3
- one nucleotide type includes a molecular tag bonded to a part of the nucleotide that is released upon 4
- monomer nucleotide incorporation, where at least one of the tags has a fluorescence property that 5
- undergoes a change before, during and/or after each of a sequence of monomer nucleotide 6 incorporations due to an interaction between the polymerizing agent polymerase tag and the
- 7 monomer nucleotide tag and where the site comprises a naturally occurring cysteine site or a 8
- cysteine replacement site in the polymerizing agent polymerase selected so that the site is less than 9
- or equal to about 50Å from a tag on each incorporating monomer nucleotide and is a site that is not 10
- involved in the function of the polymerizing agent polymerase and the polymerizing agent 11
- polymerase tag is covalently bonded to the naturally occurring cysteine site or the cysteine 12
- replacement site through its SH group. 13
 - The composition of claim 79, wherein the site is less than or equal to 80.(currently amended) 1
 - about 15Å from a tag on each incorporating monomer nucleotide. 2
 - The composition of claim 79, wherein the site is less than or equal to 81.(currently amended) 1
 - about 10Å from a tag on each incorporating monomer nucleotide. 2
 - The composition of claim 79, wherein the polymerizing agent is a 82.(currently amended) 1
 - polymerase or comprises a reverse transcriptase. 2
 - The composition of claim 79, wherein the polymerase is selected from 83.(previously presented) l

the group consisting of Taq DNA polymerase I, T7 DNA polymerase, Sequenase, and the Klenow 2 fragment from E. coli DNA polymerase I. 3 The composition of claim 82, wherein the reverse transcriptase 84.(previously presented) 1 comprises HIV-1 reverse transcriptase. 2 The composition of claim 79, wherein each of the monomers 85.(currently amended) 1 nucleotides comprises a deoxynucleotide triphosphate (dNTP) and the monomer nucleotide tag is 2 covalently bonded directly or through a linker to the pyrophosphate moiety of each its dNTP. 3 The composition of claim 8579, wherein the tag comprise fluorescent 86.(currently amended) 1 tag and the fluorescence property comprises a duration, an intensity and/or frequency of emitted 2 fluorescent light. 3 The composition of claim 8679, wherein the polymerase tag comprises 87.(currently amended) 1 a fluorescent tag and wherein the fluorescence property is fluorescence resonance energy transfer 2 (FRET), where either the monomer nucleotide tag or the polymerase tag comprises a donor and the 3 other tag comprises an acceptor and where FRET occurs when the two tags are in close proximity. 4 The composition of claim 83, wherein the polymerase comprises Taq 88.(currently amended) 5 DNA polymerase I having a tag attached to an amino acid at a specific amino acid position of the 6 Taq-DNA polymerase I, where the amino acid position and the site is selected from the group 7 consisting of 513-518, 643, 647, 649 and 653-661 of SEQ. ID No. 11, where the tag comprises a 8 9 fluorescent molecule. A composition comprising a polymerizing agent polymerase including 89.(currently amended) 1 a molecular tag covalently bonded to a site on the polymerizing agent polymerase and a monomer 2 nucleotide including a molecular tag covalently bonded to the monomer and a part of the nucleotide 3 that is released upon monomer nucleotide incorporation, where at least one of the tags has a

fluorescence property that undergoes a change before, during and/or after each of a sequence of

monomer nucleotide incorporations due to an interaction between the polymerizing agent

polymerase tag and the monomer nucleotide tag and where the site comprises a naturally occurring

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- cysteine site or a cysteine replacement site in the polymerizing agent polymerase selected so that 8 the site is less than or equal to about 50Å from a tag on each incorporating monomer nucleotide and 9 the polymerizing agent polymerase tag is covalently bonded to the naturally occurring cysteine site 10 or the cysteine replacement site through its SH group. 11
 - The composition of claim 89, wherein the site is less than or equal to 90.(currently amended) 1 about 15Å from a tag on each incorporating monomer nucleotide. 2
 - The composition of claim 89, wherein the site is less than or equal to 91.(currently amended) 1 about 10Å from a tag on each incorporating monomer nucleotide. 2
 - The composition of claim 89, wherein the polymerizing agent is a 92.(currently amended) 1 polymerase or comprises a reverse transcriptase. 2
 - 93.(canceled) 1
 - The composition of claim 9289, wherein the polymerase is selected 94.(currently amended) 1
 - from the group consisting of Taq DNA polymerase I, T7 DNA polymerase, Sequenase, and the 2
 - Klenow fragment from E. coli DNA polymerase I. 3
 - The composition of claim 92, wherein the reverse transcriptase 95.(previously presented) 1 comprises HIV-1 reverse transcriptase. 2
 - The composition of claim 89, wherein each of the monomers 96.(currently amended) 1 nucleotides comprises a deoxynucleotide triphosphate (dNTP) and the monomer nucleotide tag is 2
 - covalently bonded directly or through a linker to the terminal phosphate group of each its dNTP. 3
 - The composition of claim 9689, wherein the tags comprise fluorescent 97.(currently amended) 1
 - tags and the fluorescence property comprises a duration, an intensity and/or frequency of emitted 2
 - fluorescent light. 3
 - The composition of claim 97, wherein the polymerase tag comprises 98.(currently amended) 1
 - a fluorescent tag and wherein the fluorescence property is fluorescence resonance energy transfer 2

- (FRET), where either the monomer nucleotide tag or the polymerase tag comprises a donor and the 3 other tag comprises an acceptor and where FRET occurs when the two tags are in close proximity. 4
- The composition of claim 94, wherein the polymerase comprises Taq 99.(previously presented) 5
- DNA polymerase I having a tag attached to an amino acid at a specific amino acid position of the 6
- Taq DNA polymerase I, where the amino acid position and the site is selected from the group 7
- consisting of 513-518, 643, 647, 649 and 653-661 of SEQ. ID No. 11, where the tag comprises a 8
- fluorescent molecule. 9
- 100.(previously presented) The composition of claim 50, wherein the polymerizing agent lacks 1
- the ability to remove a previously incorporated monomer nucleotide. 2
 - 101.(canceled)
- 102.(previously presented) The composition of claim 64, wherein the polymerase is free of or 1
- lacks the ability to remove a previously incorporated monomer nucleotide. 2
- 103.(previously presented) The composition of claim 71, wherein the polymerase is free of or 1
- lacks the ability to remove a previously incorporated monomer nucleotide. 2
- 104.(previously presented) The composition of claim 89, wherein the polymerase is free of or 1
- lacks the ability to remove a previously incorporated monomer nucleotide. 2
- 105.(previously presented) The composition of claim 79, wherein the site is less than or equal to 1
- about 25Å from a tag on each incorporating monomer nucleotide. 2
- 106.(previously presented) The composition of claim 89, wherein the site is less than or equal to 1
- about 25Å from a tag on each incorporating monomer nucleotide. 2
 - 107.(canceled)
- The composition of claim 10, wherein the polymerase comprises a genetically 108.(new) 1
- engineered polymerase comprising a native polymerase including one cysteine residue replacement 2
- or a plurality of cysteine residue replacements at one site or a plurality of sites of the native 3

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polymerase, where the site or sites are not in contact with other proteins, where the site or sites do not alter the conformation or folding of the polymerase, where the site or sites are not involved in the functioning of the polymerase, and where the polymerase tag is bonded to the polymerase through a cysteine residue replacement or through a plurality of cysteine residue replacements.

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The composition of claim 50, wherein the polymerase comprises a genetically 109.(new) engineered polymerase comprising a native polymerase including one cysteine residue replacement or a plurality of cysteine residue replacements at one site or a plurality of sites of the native polymerase, where the site or sites are not in contact with other proteins, where the site or sites do not alter the conformation or folding of the polymerase, where the site or sites are not involved in the functioning of the polymerase, and where the polymerase tag is bonded to the polymerase through a cysteine residue replacement or through a plurality of cysteine residue replacements.

The composition of claim 64, wherein the polymerase comprises a genetically 110.(new) engineered polymerase comprising a native polymerase including one cysteine residue replacement or a plurality of cysteine residue replacements at one site or a plurality of sites of the native polymerase, where the site or sites are not in contact with other proteins, where the site or sites do not alter the conformation or folding of the polymerase, where the site or sites are not involved in the functioning of the polymerase, and where the polymerase tag is bonded to the polymerase through a cysteine residue replacement or through a plurality of cysteine residue replacements.

The composition of claim 71, wherein the polymerase comprises a genetically 111.(new) engineered polymerase comprising a native polymerase including one cysteine residue replacement or a plurality of cysteine residue replacements at one site or a plurality of sites of the native polymerase, where the site or sites are not in contact with other proteins, where the site or sites do not alter the conformation or folding of the polymerase, where the site or sites are not involved in the functioning of the polymerase, and where the polymerase tag is bonded to the polymerase through a cysteine residue replacement or through a plurality of cysteine residue replacements.